SEQUENCE LISTING

<110> Bistrup, Annette
Rosen, Steven D.
Tangemann, Kirsten
Hemmerich, Stefan

<120> GLYCOSYL SULFOTRANSFERASE-3

<130> UCAL-107DIV

<150> 09/190,911
<151> 1998-11-12

<150> 09/045,284
<151> 1998-03-20

<160> 8

<170> FastSEQ for Windows Version 3.0

<210> 1
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<213> H. sapiens

<400> 1

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            20                  25                  30
Ser Leu Ser Met Lys Ala Gln Pro Glu Arg Met His Val Leu Val Leu
        35                  40                  45
Ser Ser Trp Arg Ser Gly Ser Ser Phe Val Gly Gln Leu Phe Gly Gln
    50                  55                  60
His Pro Asp Val Phe Tyr Leu Met Glu Pro Ala Trp His Val Trp Met
65                  70                  75                  80
Thr Phe Lys Gln Ser Thr Ala Trp Met Leu His Met Ala Val Arg Asp
                85                  90                  95
Leu Ile Arg Ala Val Phe Leu Cys Asp Met Ser Val Phe Asp Ala Tyr
            100                 105                 110
Met Glu Pro Gly Pro Arg Arg Gln Ser Ser Leu Phe Gln Trp Glu Asn
        115                 120                 125
Ser Arg Ala Leu Cys Ser Ala Pro Ala Cys Asp Ile Ile Pro Gln Asp
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Glu Ile Ile Pro Arg Ala His Cys Arg Leu Leu Cys Ser Gln Gln Pro
145                 150                 155                 160
Phe Glu Val Val Glu Lys Ala Cys Arg Ser Tyr Ser His Val Val Leu
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Lys Glu Val Arg Phe Phe Asn Leu Gln Ser Leu Tyr Pro Leu Leu Lys
            180                 185                 190
Asp Pro Ser Leu Asn Leu His Ile Val His Leu Val Arg Asp Pro Arg
        195                 200                 205
Ala Val Phe Arg Ser Arg Glu Arg Thr Lys Gly Asp Leu Met Ile Asp
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Ser Arg Ile Val Met Gly Gln His Glu Gln Lys Leu Lys Lys Glu Asp
225                 230                 235                 240
Gln Pro Tyr Tyr Val Met Gln Val Ile Cys Gln Ser Gln Leu Glu Ile
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Tyr Lys Thr Ile Gln Ser Leu Pro Lys Ala Leu Gln Glu Arg Tyr Leu
            260                 265                 270
Leu Val Arg Tyr Glu Asp Leu Ala Arg Ala Pro Val Ala Gln Thr Ser
        275                 280                 285
Arg Met Tyr Glu Phe Val Gly Leu Glu Phe Leu Pro His Leu Gln Thr
    290                 295                 300
Trp Val His Asn Ile Thr Arg Gly Lys Gly Met Gly Asp His Ala Phe
305                 310                 315                 320
His Thr Asn Ala Arg Asp Ala Leu Asn Val Ser Gln Ala Trp Arg Trp
                325                 330                 335
Ser Leu Pro Tyr Glu Lys Val Ser Arg Leu Gln Lys Ala Cys Gly Asp
            340                 345                 350
Ala Met Asn Leu Leu Gly Tyr Arg His Val Arg Ser Glu Gln Glu Gln
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Arg Asn Leu Leu Leu Asp Leu Leu Ser Thr Trp Thr Val Pro Glu Gln
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Ile His
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<220>
<223> primer

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<210> 4
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<221> misc_feature
<222> (1)...(29)
<223> n = inosine

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<210> 5
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<221> misc_feature
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<223> n = inosine

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<210> 6
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<221> misc_feature
<222> (1)...(26)
<223> n = inosine

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<211> 37
<212> DNA
<213> H. sapiens

<400> 7
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<210> 8
<211> 47
<212> DNA
<213> H. sapiens

<400> 8
ataaagcttg tggatttgtt cagggacatt ccaggtagac agaagat 47